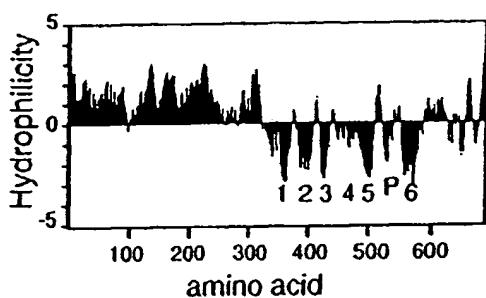


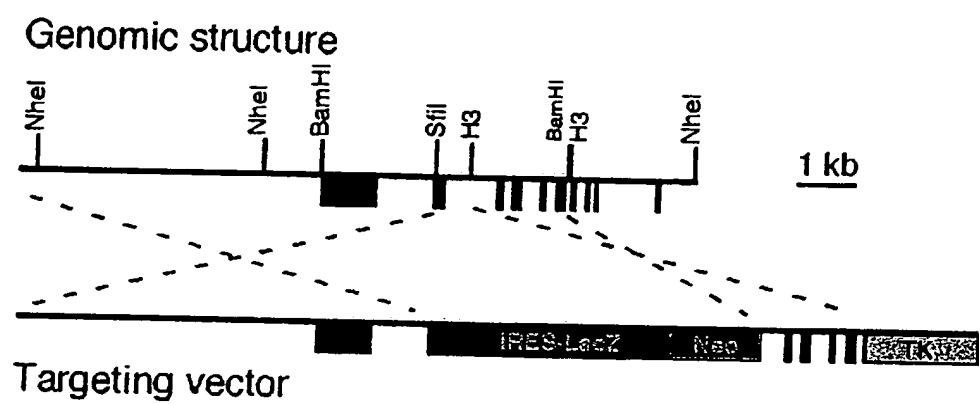
**a**

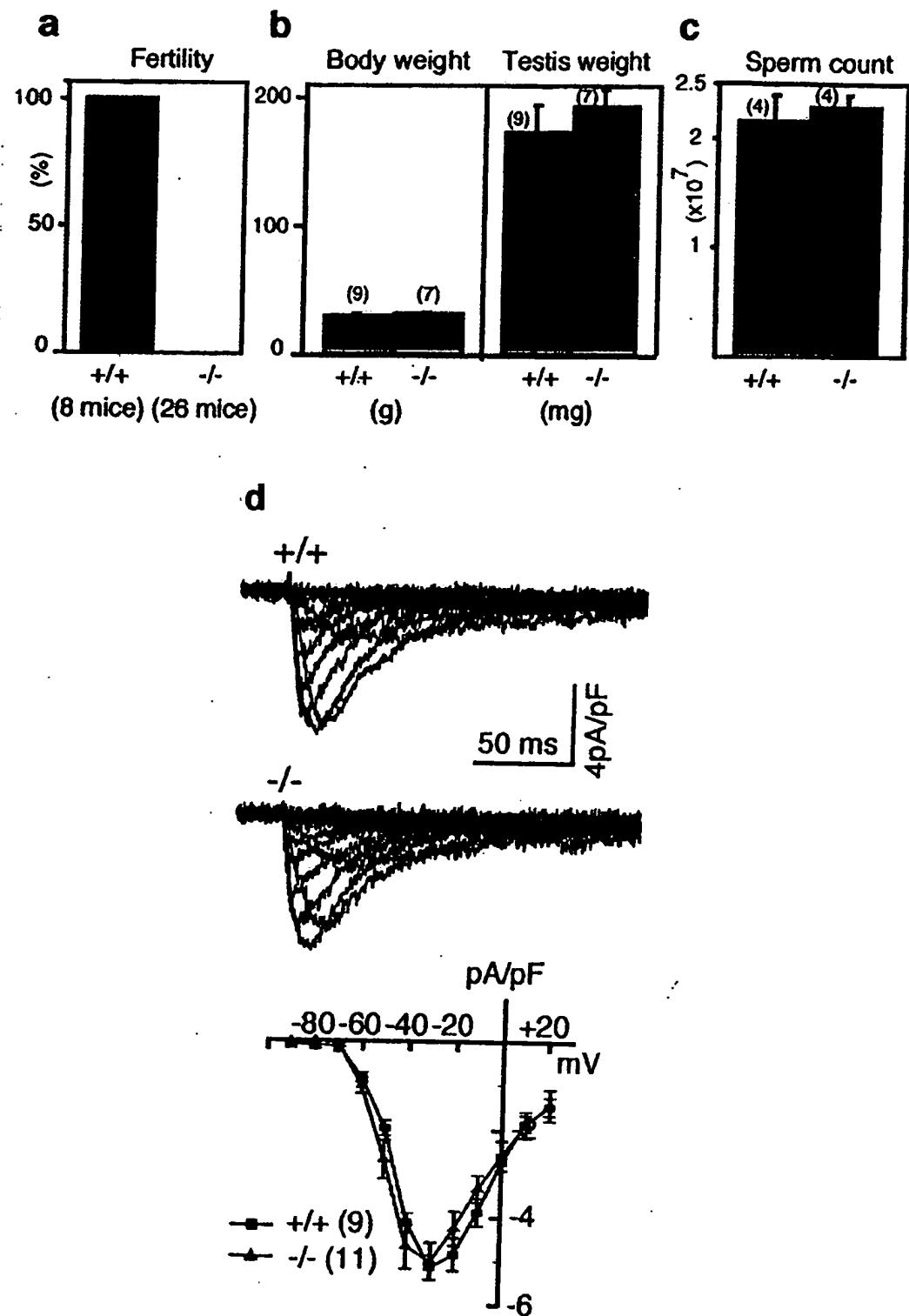
MDQSSRRDESYHETHPGSLDPHQSHPHPHPHPTLHRPNQGGVYDSPQH 50  
 GMFQQPYQQHGGFHQQNELQHLREFSDSHDNAFSHHSYQQDRAGVSTLPN 100  
 NISHAYGGSHPLAESQHSGGPQSGPRIDPNHHPHQDDPHRSEPLSHPSS 150  
 TGSHQGTTHQHQYHERSHHLNPQQNRDHADTISYRSSTRFYRSHAPFSRQE 200  
 RPHLHADHHHEGHHAHSHHGEHPHKEQQRHYHGDHMHHIHRSPSASQL 250  
 SEGEDHVQKRKKAQRAHKKAHGTGNIFQLLWEKISHLLLGLQQMILSLTQS 350  
 S1   S2  
 LGFETFIFIVVCLNTVILVAQT~~FTELEIRGEWYF~~**MVLDSIFLSIYVLEAV** 400  
 S3  
 LKLIAL~~GLEYFYDPWNNLDFFIMVM~~**AVLDVFULLQINSLSYSFYNHSI**~~FRI~~ 450  
 S4  
 LKVFKSMRALRAIRVLRR~~I~~**SILTSLHEVAGTLSGS**~~LPSITAILTL~~**MFTCL** 500  
 P  
 S6  
 FLFSVVLRA~~Q~~**FQDSDPKRFQN**~~IFTTLFTLFTMLTLDDWSL~~**IYIDNRAQGA** 550  
 WYIIPILMIVIVI~~OYFIFLN~~**LNVIAVLVDNFQMALLKG**LEKVVKLEQAARVH 600  
 EKLLDDSLTDLNKADANAQMTEEALKMQLIEGMFGNMTVKQRVLHFQFLQ 650  
 LVAAVEQHQHQKFRSQAYVIDELVDMAFEAGDDDYGK 686

**b****c****Pore region**

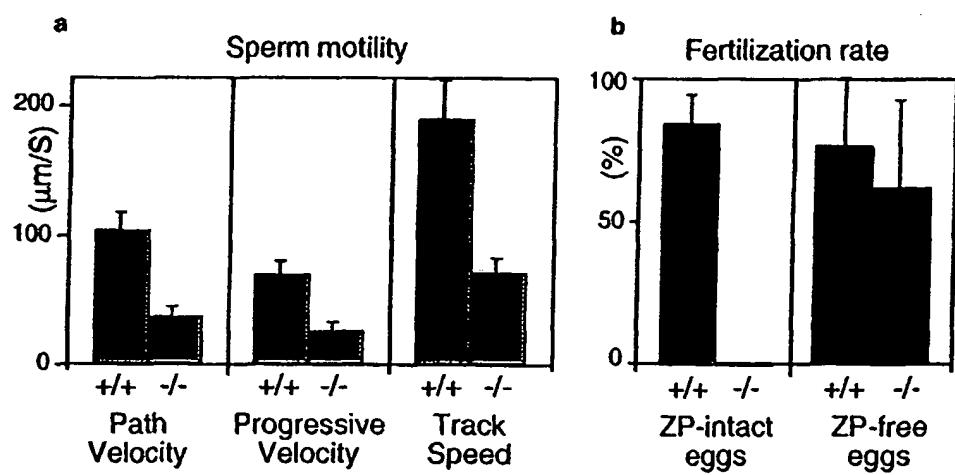
CatSper	RFQNI <del>FTT</del> <b>LFTLFTMLI</b> QD <del>NS</del> <b>SIYID</b>
Cav1.2	NFDNF <del>AF</del> <b>AMLTVF</b> QCITMEGWT <del>D</del> VLYN
I Cav2.2	NFDNL <del>I</del> <b>FAILTVF</b> QCITMEGWT <del>D</del> ILYN
Cav3.1	NFDNIGYAWIAIFQV <del>ITL</del> CG/DIMYF
Cav1.2	TFDNFPQ <del>SL</del> <b>LTVF</b> QILIGED <del>D</del> NSVMD
II Cav2.2	NFDTPA <del>AI</del> <b>LTVF</b> QILIGED <del>D</del> NAVMYN
Cav3.1	NFD <del>S</del> LLWA <del>I</del> <b>VT</b> VFQILIGED <del>D</del> QNKLYN
Cav1.2	DFDNVLAAMMALFTV <del>ST</del> CGPPELLYR
III Cav2.2	HYDNVLW <del>ALL</del> <b>LT</b> FTV <del>ST</del> CGGQPMYLKH
Cav3.1	NFDNLGQALMSLFV <del>LA</del> SK <del>QG</del> /DIMYD
Cav1.2	NFQTFPQAV <del>LL</del> <b>PRCAT</b> GEA <del>Q</del> DIMLA
IV Cav2.2	NFR <del>T</del> FLQALM <del>LL</del> <b>FRSA</b> GEA <del>Q</del> HEIMLS
Cav3.1	TFRNFGMA <del>FL</del> <b>TL</b> FRV <del>ST</del> CG <del>Q</del> NGIMKD

**FIG. 1**

**FIG. 2**



**FIG. 3**

**FIG. 4**

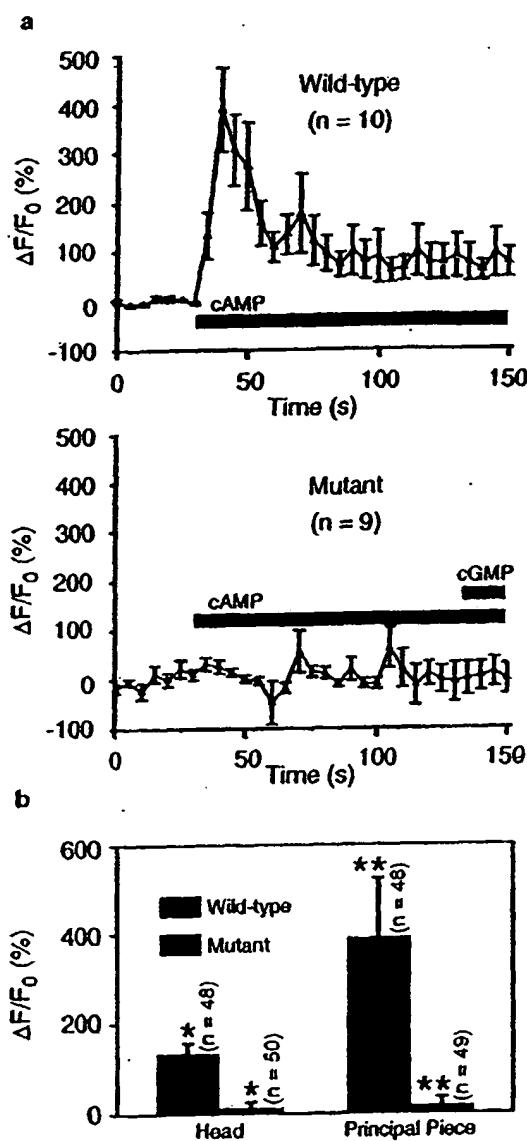


FIG. 5